	CRF Errors Corrected by the STIC Systems Branch	/600 Dat : 11/1/200
mber: Changed	1 a file from non-ASCII to ASCII	(STIC st
hanged	the margins in cases where the sequence text was "wrapped" down to the ne	ext line.
-	format error in the Current Application Data section, specifically:	RECEIV
dited the	e Current Application Data section with the actual current number. The number was the prior application data; or the other	NOV 1 2 2 er inputted by the TECH CENTER 16
	e mandatory heading and subheadings for "Current Application Data".	
dited the	e "Number of Sequences" field. The applicant spelled out a number instead o	of using an integer.
hanged	the spelling of a mandatory field (the headings or subheadings), specifically:	
orrected	the SEQ ID NO when obviously incorrect. The sequence numbers that were	edited were:
serted o	or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edite	rd:
orrected oplicant	I subheading placement. All responses must be on the same line as each subplaced a response below the subheading, this was moved to its appropriate p	oheading. If the
nserted	colons after headings/subheadings. Headings edited included:	
Deleted a	extra, invalid, headings used by an applicant, specifically:	
Deleted:	non-ASCII "garbage" at the beginning/end of files; secretary initials/fe numbers throughout text; other invalid text, such as	ilename at end of fil
nserted	mandatory headings, specifically:	
Correcte	d an obvious error in the response, specifically:	11.14
Edited id	lentifiers where upper case is used but lower case is required, or vice versa.	
Correcte	d an error in the Number of Sequences field, specifically:	
\ "Hard	Page Break" code was inserted by the applicant. All occurrences had to be de	eleted.
eleted <i>e</i> ue to a P	ndIng stop codon in amino acid sequences and adjusted the "(A)Length:" field attention bug). Sequences corrected:	d accordingly (error
Other:		

*Examiner: The abov corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



1600

RAW SEQUENCE LISTING DATE: 11/07/2002 PATENT APPLICATION: US/09/844,861A TIME: 20:57:29

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\11072002\I844861A.raw

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3 <110> APPLICANT: Padigaru, Muralidhara
         Mishra, Vishnu
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 5
         Spytek, Kimberly
         Burgess, Catherine
 7
         Lepley, Denise
         Grosse, William
 8
         Szekeres, Edward
 9
10
         Alsobrook, John
11
         Gangolli, Esha
12
         Casman, Stacie
13
         MacDougall, John
         Smithson, Glennda
16 <120> TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
18 <130> FILE REFERENCE: 15966-789 US
20 <140> CURRENT APPLICATION NUMBER: 09/844,861A
21 <141> CURRENT FILING DATE: 2001-04-27
23 <150> PRIOR APPLICATION NUMBER: 60/199,947
24 <151> PRIOR FILING DATE: 2000-04-27
26 <150> PRIOR APPLICATION NUMBER: 60/199,960
27 <151> PRIOR FILING DATE: 2000-04-27
29 <150> PRIOR APPLICATION NUMBER: 60/225,226
30 <151> PRIOR FILING DATE: 2000-08-14
32 <150> PRIOR APPLICATION NUMBER: 60/256,399
33 <151> PRIOR FILING DATE: 2000-12-18
35 <150> PRIOR APPLICATION NUMBER: 60/256,524
36 <151> PRIOR FILING DATE: 2000-12-18
38 <150> PRIOR APPLICATION NUMBER: 60/258,159
39 <151> PRIOR FILING DATE: 2000-12-22
41 <150> PRIOR APPLICATION NUMBER: 60/258,511
42 <151> PRIOR FILING DATE: 2000-12-28
44 <150> PRIOR APPLICATION NUMBER: 60/258,828
45 <151> PRIOR FILING DATE: 2000-12-28
47 <150> PRIOR APPLICATION NUMBER: 60/259,659
48 <151> PRIOR FILING DATE: 2001-01-04
50 <150> PRIOR APPLICATION NUMBER: 60/275,604
51 <151> PRIOR FILING DATE: 2001-03-13
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58 <211> LENGTH: 1016
59 <212> TYPE: DNA
60 <213> ORGANISM: Homo sapiens
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65 agocottota gggaacatca ttatootgtt tgtgatacag actgaacaga gcotocacca 180
66 acceatgttt tactteetag ceatgttgge eggeactgat etgggettgt etacageaac 240
67 catccccaag atgctgggaa ttttctggtt taatcttgga gagattgcat ttggtgcctg 300
68 catcacacag atgtatacca ttcatatatg cactggcctg gagtctgtgg tactgacagt 360
69 cacgggcata gatcgctata ttgccatctg caaccccctg agatatagca tgatccttac 420
70 caacaaggta atagccattc tgggcatagt catcattgtc aggactttgg tatttgtgac 480
71 tecatteaca ttteteacee tgagattgee tttetgtggt gteeggatta teceteatae 540
72 ctattgtgaa cacatgggct tggcaaagtt agcttgtgcc agtattaatg ttatatatgg 600
73 attgattgcc ttctcagtgg gatacattga catttctgtg attggatttt cctatgtcca 660
74 gatecteega getgtettee ateteecage etgggatgee eggettaagg caeteageae 720
75 atgtggctct cacgtctgtg ttatgttggc tttctacctg ccagccctct tttccttcat 780
76 gacacaccgc tttggccaca acatccctca ttacatccac attcttctgg ccaatctgta 840
77 tgtggttttt cccctgctc ttaactctgt tatctatggg gtcaaaacaa aacagatacg 900
78 agagcaggta cttaggatac tcaaccctaa aagcttttgg cattttgacc ccaagaggat 960
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91 Val Gly Val Pro Gly Leu Glu Asp Val His Val Trp Ile Gly Phe Pro
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94 Phe Phe Ala Val Tyr Leu Thr Ala Leu Leu Gly Asn Ile Ile Ile Leu
                                40
97 Phe Val Ile Gln Thr Glu Gln Ser Leu His Gln Pro Met Phe Tyr Phe
                            55
100 Leu Ala Met Leu Ala Gly Thr Asp Leu Gly Leu Ser Thr Ala Thr Ile
101
                                              75
103 Pro Lys Met Leu Gly Ile Phe Trp Phe Asn Leu Gly Glu Ile Ala Phe
104
                     85
                                         90
106 Gly Ala Cys Ile Thr Gln Met Tyr Thr Ile His Ile Cys Thr Gly Leu
107
                100
                                    105
109 Glu Ser Val Val Leu Thr Val Thr Gly Ile Asp Arg Tyr Ile Ala Ile
110
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                                                     125
112 Cys Asn Pro Leu Arg Tyr Ser Met Ile Leu Thr Asn Lys Val Ile Ala
113
        130
                            135
                                                 140
115 Ile Leu Gly Ile Val Ile Ile Val Arg Thr Leu Val Phe Val Thr Pro
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                                            155
118 Phe Thr Phe Leu Thr Leu Arg Leu Pro Phe Cys Gly Val Arg Ile Ile
119
                                        170
                    165
121 Pro His Thr Tyr Cys Glu His Met Gly Leu Ala Lys Leu Ala Cys Ala
122
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124 Ser Ile Asn Val Ile Tyr Gly Leu Ile Ala Phe Ser Val Gly Tyr Ile
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127 Asp Ile Ser Val Ile Gly Phe Ser Tyr Val Gln Ile Leu Arg Ala Val
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Input Set : A:\PTO.AMC.txt

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133 Gly Ser His Val Cys Val Met Leu Ala Phe Tyr Leu Pro Ala Leu Phe
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136 Ser Phe Met Thr His Arg Phe Gly His Asn Ile Pro His Tyr Ile His
137
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                                    265
                                                         270
139 Ile Leu Leu Ala Asn Leu Tyr Val Val Phe Pro Pro Ala Leu Asn Ser
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142 Val Ile Tyr Gly Val Lys Thr Lys Gln Ile Arg Glu Gln Val Leu Arg
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154 <212> TYPE: DNA
155 <213> ORGANISM: Homo sapiens
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159 tqtcaqccat catqtctqqq qacaacaqct ccaqcctqac cccaqqattc tttatcttqa 120
160 atggcgttcc tgggctggaa gccacacaca tctggatctc cctgccattc tgctttatgt 180
161 acatcattgc tgtcgtgggg aactgtgggc tcatctgcct catcagccat gaggaggccc 240
162 tgcaccggcc catgtactac ttcctggccc tgctctcctt cactgatgtc accttgtgca 300
163 ccaccatggt acctaatatg ctgtgcatat tctggttcaa cctcaaggag attgacttta 360
164 acgcctgcct ggcccagatg ttttttgtcc atatgctgac agggatggag tctggggtgc 420
165 tcatgctcat ggccctggac cgctatgtgg ccatctgcta ccccttacgc tatgccacca 480
166 teettaceaa eeetgteate geeaaggetg gtettgeeae ettettgagg aatgtgatge 540
167 tcatcatccc attcactctc ctcaccaagc gcctgcccta ttgccggggg aacttcatcc 600
168 cccacaccta ctgtgaccat atgtctgtgg ccaaggtatc ctgtggcaat ttcaaggtca 660
169 atgctattta tggtctgatg gttgctctcc tgattggtgt gtttgatatc tgctgtatct 720
170 ctgtatctta cactatgatt ttgcaggctg ttatgagcct gtcatcagca gatgctcgtc 780
171 acaaageett cagcacetge acateteaca tgtgtteeat tgtgateace tatgttgetg 840
172 cttttttcac ttttttcact catcgttttg taggacacaa tatcccaaac cacatacaca 900
173 tcatcgtggc caacctttat ctgctactgc ctcctaccat gaacccaatt gtttatggag 960
174 tcaagaccaa gcagattcag gaaggtgtaa ttaaattttt acttggagac aaggttagtt 1020
175 ttacctatga caaatgaaac atagaataga catattgttt caggtggtga gaaaataatg 1080
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180 <211> LENGTH: 336
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191 Gly Val Pro Gly Leu Glu Ala Thr His Ile Trp Ile Ser Leu Pro Phe
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Input Set : A:\PTO.AMC.txt

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200 Ala Leu Leu Ser Phe Thr Asp Val Thr Leu Cys Thr Thr Met Val Pro
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                                         90
203 Asn Met Leu Cys Ile Phe Trp Phe Asn Leu Lys Glu Ile Asp Phe Asn
                100
                                    105
206 Ala Cys Leu Ala Gln Met Phe Phe Val His Met Leu Thr Gly Met Glu
207
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                                120
209 Ser Gly Val Leu Met Leu Met Ala Leu Asp Arg Tyr Val Ala Ile Cys
        130
                            135
                                                 140
212 Tyr Pro Leu Arg Tyr Ala Thr Ile Leu Thr Asn Pro Val Ile Ala Lys
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                                            155
215 Ala Gly Leu Ala Thr Phe Leu Arg Asn Val Met Leu Ile Ile Pro Phe
216
                    165
                                        170 ·
218 Thr Leu Leu Thr Lys Arg Leu Pro Tyr Cys Arg Gly Asn Phe Ile Pro
219
                180
                                    185
                                                         190
221 His Thr Tyr Cys Asp His Met Ser Val Ala Lys Val Ser Cys Gly Asn
            195
                                200
                                                     205
224 Phe Lys Val Asn Ala Ile Tyr Gly Leu Met Val Ala Leu Leu Ile Gly
                            215
227 Val Phe Asp Ile Cys Cys Ile Ser Val Ser Tyr Thr Met Ile Leu Gln
                        230
                                            235
230 Ala Val Met Ser Leu Ser Ser Ala Asp Ala Arg His Lys Ala Phe Ser
231
                    245
                                        250
233 Thr Cys Thr Ser His Met Cys Ser Ile Val Ile Thr Tyr Val Ala Ala
                                    265
                260
236 Phe Phe Thr Phe Phe Thr His Arg Phe Val Gly His Asn Ile Pro Asn
                                                     285
237
            275
                                280
239 His Ile His Ile Ile Val Ala Asn Leu Tyr Leu Leu Pro Pro Thr
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                            295
                                                 300
242 Met Asn Pro Ile Val Tyr Gly Val Lys Thr Lys Gln Ile Gln Glu Gly
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260 ttggaagatg tgcatttgtg gatctccttc ccactgtgta ccatgtacag cattgctatt 180
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262 tatgtettee ttgecettet tteetteaca gatgtgetea tgtgeaceag caecetteee 300
263 aacactctct tcatattgtg gtttaatctc aaggagattg attttaaagc ctgcctcgcc 360
264 cagatgttct ttgtgcacac cttcacaggg atggagtctg gggtgctcat gctcatggcc 420
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Input Set : A:\PTO.AMC.txt

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266 gtcattgcta aagctgggtt cctcactttt cttaggggtg tgatgcttgt tatcccttcc 540
267 actttcctca ccaagegeet tecatactge aagggeaacg teataceeca cacetactgt 600
268 gaccacatgt ctgtggccaa gatatcttgt ggtaatgtca gggttaacgc catctatggt 660
269 ttgatagttg ccctgctgat tgggggcttt gatatcctgt gcattacaat ctcctacact 720
270 atgattette aageagttgt gagtetatea teageagatg etegacagaa ggeetteage 780
271 acctgcactg cocacttctg tgccatagtc ctcacctatg ttccagcctt ctttaccttc 840
272 tttacacacc attttggggg acacaccatt cctctacaca tacatattat tatggctaat 900
273 ctctacctac taatgcctcc cacaatgaac cctattgtgt atggggtgaa aaccaggcag 960
274 gtacgagaaa gtgtcattag gttctttctt aagggaaagg acaattctca taacttttaa 1020
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                20
290 Leu Cys Thr Met Tyr Ser Ile Ala Ile Thr Gly Asn Phe Gly Leu Met
                                 40
293 Tyr Leu Ile Tyr Cys Asp Glu Ala Leu His Arg Pro Met Tyr Val Phe
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                             55
296 Leu Ala Leu Leu Ser Phe Thr Asp Val Leu Met Cys Thr Ser Thr Leu
297 65
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299 Pro Asn Thr Leu Phe Ile Leu Trp Phe Asn Leu Lys Glu Ile Asp Phe
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302 Lys Ala Cys Leu Ala Gln Met Phe Phe Val His Thr Phe Thr Gly Met
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305 Glu Ser Gly Val Leu Met Leu Met Ala Leu Asp His Cys Val Ala Ile
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308 Cys Phe Pro Leu Arg Tyr Ala Thr Ile Leu Thr Asn Ser Val Ile Ala
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311 Lys Ala Gly Phe Leu Thr Phe Leu Arg Gly Val Met Leu Val Ile Pro
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314 Ser Thr Phe Leu Thr Lys Arg Leu Pro Tyr Cys Lys Gly Asn Val Ile
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315
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317 Pro His Thr Tyr Cys Asp His Met Ser Val Ala Lys Ile Ser Cys Gly
318
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320 Asn Val Arg Val Asn Ala Ile Tyr Gly Leu Ile Val Ala Leu Leu Ile
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                                                     205
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                                                 220
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327 225
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329 Ser Thr Cys Thr Ala His Phe Cys Ala Ile Val Leu Thr Tyr Val Pro
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VERIFICATION SUMMARY

DATE: 11/07/2002

PATENT APPLICATION: US/09/844,861A

TIME: 20:57:30

Input Set : A:\PTO.AMC.txt